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RAW SEQUENCE LISTING DATE: 03/05/2002 P.5
PATENT APPLICATION: US/10/078,059 TIME: 14:12:03

Input Set : A:\PF466P2SEQLIST01102002.txt
Output Set: N:\CRF3\03052002\J078059.raw

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      3 <110> APPLICANT: Ruben et al.
      5 <120> TITLE OF INVENTION: Cytokine Receptor Common Gamma Chain Like
      7 <130> FILE REFERENCE: PF466P2
C-->  9 <140> CURRENT APPLICATION NUMBER: US/10/078,059
C-->  9 <141> CURRENT FILING DATE: 2002-02-20
      9 <150> PRIOR APPLICATION NUMBER: 60/269,876
     10 <151> PRIOR FILING DATE: 2001-02-21
     12 <150> PRIOR APPLICATION NUMBER: PCT/US00/22493
     13 <151> PRIOR FILING DATE: 2000-08-17
     15 <150> PRIOR APPLICATION NUMBER: 09/376,430
     16 <151> PRIOR FILING DATE: 1999-08-18
     18 <150> PRIOR APPLICATION NUMBER: 09/263,626
     19 <151> PRIOR FILING DATE: 1999-03-05
     21 <150> PRIOR APPLICATION NUMBER: PCT/US99/05068
     22 <151> PRIOR FILING DATE: 1999-03-05
     24 <150> PRIOR APPLICATION NUMBER: 60/086,505
     25 <151> PRIOR FILING DATE: 1998-05-22
     27 <150> PRIOR APPLICATION NUMBER: 60/078,563
     28 <151> PRIOR FILING DATE: 1998-03-19
     30 <160> NUMBER OF SEQ ID NOS: 32
     32 <170> SOFTWARE: PatentIn Ver. 2.1
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     35 <211> LENGTH: 1573
     36 <212> TYPE: DNA
     37 <213> ORGANISM: Homo sapiens
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     40 <221> NAME/KEY: CDS
     41 <222> LOCATION: (13)..(1125)
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     48 ctg ctg gga ggc tgg atg gct ttg ggg caa gga gga gca gca gaa gga   99
     49 Leu Leu Gly Gly Trp Met Ala Leu Gly Gln Gly Gly Ala Ala Glu Gly
     50      15                  20                  25
     52 gta cag att cag atc atc tac ttc aat tta gaa acc gtg cag gtg aca   147
     53 Val Gln Ile Gln Ile Ile Tyr Phe Asn Leu Glu Thr Val Gln Val Thr
     54  30                  35                  40                  45
     56 tgg aat gcc agc aaa tac tcc agg acc aac ctg act ttc cac tac aga   195
     57 Trp Asn Ala Ser Lys Tyr Ser Arg Thr Asn Leu Thr Phe His Tyr Arg
     58                  50                  55                  60
     60 ttc aac ggt gat gag gcc tat gac cag tgc acc aac tac ctt ctc cag   243
     61 Phe Asn Gly Asp Glu Ala Tyr Asp Gln Cys Thr Asn Tyr Leu Leu Gln
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Input Set : A:\PF466P2SEQLIST01102002.txt
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64 gaa ggt cac act tcg ggg tgc ctc cta gac gca gag cag cga gac gac   291
65 Glu Gly His Thr Ser Gly Cys Leu Leu Asp Ala Glu Gln Arg Asp Asp
66           80                  85                  90
68 att ctc tat ttc tcc atc agg aat ggg acg cac ccc gtt ttc acc gca   339
69 Ile Leu Tyr Phe Ser Ile Arg Asn Gly Thr His Pro Val Phe Thr Ala
70       95                 100                 105
72 agt cgc tgg atg gtt tat tac ctg aaa ccc agt tcc ccg aag cac gtg   387
73 Ser Arg Trp Met Val Tyr Tyr Leu Lys Pro Ser Ser Pro Lys His Val
74 110                  115                 120                 125
76 aga ttt tcg tgg cat cag gat gca gtg acg gtg acg tgt tct gac ctg   435
77 Arg Phe Ser Trp His Gln Asp Ala Val Thr Val Thr Cys Ser Asp Leu
78                  130                 135                 140
80 tcc tac ggg gat ctc ctc tat gag gtt cag tac cgg agc ccc ttc gac   483
81 Ser Tyr Gly Asp Leu Leu Tyr Glu Val Gln Tyr Arg Ser Pro Phe Asp
82              145                 150                 155
84 acc gag tgg cag tcc aaa cag gaa aat acc tgc aac gtc acc ata gaa   531
85 Thr Glu Trp Gln Ser Lys Gln Glu Asn Thr Cys Asn Val Thr Ile Glu
86          160                 165                 170
88 ggc ttg gat gcc gag aag tgt tac tct ttc tgg gtc agg gtg aag gct   579
89 Gly Leu Asp Ala Glu Lys Cys Tyr Ser Phe Trp Val Arg Val Lys Ala
90      175                 180                 185
92 atg gag gat gta tat ggg cca gac aca tac cca agc gac tgg tca gag   627
93 Met Glu Asp Val Tyr Gly Pro Asp Thr Tyr Pro Ser Asp Trp Ser Glu
94 190                  195                 200                 205
96 gtg aca tgc tgg cag aga ggc gag att cgg gat gcc tgt gca gag aca   675
97 Val Thr Cys Trp Gln Arg Gly Glu Ile Arg Asp Ala Cys Ala Glu Thr
98                  210                 215                 220
100 cca acg cct ccc aaa cca aag ctg tcc aaa ttt att tta att tcc agc   723
101 Pro Thr Pro Pro Lys Pro Lys Leu Ser Lys Phe Ile Leu Ile Ser Ser
102             225                 230                 235
104 ctg gcc atc ctt ctg atg gtg tct ctc ctc ctt ctg tct tta tgg aaa   771
105 Leu Ala Ile Leu Leu Met Val Ser Leu Leu Leu Leu Ser Leu Trp Lys
106         240                 245                 250
108 tta tgg aga gtg aag aag ttt ctc att ccc agc gtg cca gac ccg aaa   819
109 Leu Trp Arg Val Lys Lys Phe Leu Ile Pro Ser Val Pro Asp Pro Lys
110     255                 260                 265
112 tcc atc ttc ccc ggg ctc ttt gag ata cac caa ggg aac ttc cag gag   867
113 Ser Ile Phe Pro Gly Leu Phe Glu Ile His Gln Gly Asn Phe Gln Glu
114 270                 275                 280                 285
116 tgg atc aca gac acc cag aac gtg gcc cac ctc cac aag atg gca ggt   915
117 Trp Ile Thr Asp Thr Gln Asn Val Ala His Leu His Lys Met Ala Gly
118                 290                 295                 300
120 gca gag caa gaa agt ggc ccc gag gag ccc ctg gta gtc cag ttg gcc   963
121 Ala Glu Gln Glu Ser Gly Pro Glu Glu Pro Leu Val Val Gln Leu Ala
122             305                 310                 315
124 aag act gaa gcc gag tct ccc agg atg ctg gac cca cag acc gag gag   1011
125 Lys Thr Glu Ala Glu Ser Pro Arg Met Leu Asp Pro Gln Thr Glu Glu
126         320                 325                 330
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Input Set : A:\PF466P2SEQLIST01102002.txt
Output Set: N:\CRF3\03052002\J078059.raw

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128 aaa gag gcc tct ggg gga tcc ctc cag ctt ccc cac cag ccc ctc caa   1059
129 Lys Glu Ala Ser Gly Gly Ser Leu Gln Leu Pro His Gln Pro Leu Gln
130     335                 340                 345
132 ggc ggt gat gtg gtc aca atc ggg ggc ttc acc ttt gtg atg aat gac   1107
133 Gly Gly Asp Val Val Thr Ile Gly Gly Phe Thr Phe Val Met Asn Asp
134 350                 355                 360                 365
136 cgc tcc tac gtg gcg ttg tgatggacac accactgtca aagtcaacgt          1155
137 Arg Ser Tyr Val Ala Leu
138                 370
140 caggatcoac gttgacattt aaagacagag gggactgtcc cggggactcc acaccaccat 1215
142 ggatgggaag tctccacgcc aatgatggta ggactaggag actctgaaga cccagcctca 1275
144 ccgcctaatg cggccactgc cctgctaact ttcccccaca tgagtctctg tgttcaaagg 1335
146 cttgatggca gatgggagcc aattgctcca ggagatttac tcccagttcc ttttcgtgcc 1395
148 tgaacgttgt cacataaacc ccaaggcagc acgtccaaaa tgctgtaaaa ccatcttccc 1455
150 actctgtgag tccccagttc cgtccatgta cctgttccat agcattggat tctcggagga 1515
152 ttttttgtct gttttgagac tccaaaccac ctctacccct acaaaaaaaa aaaaaaaa   1573
155 <210> SEQ ID NO: 2
156 <211> LENGTH: 371
157 <212> TYPE: PRT
158 <213> ORGANISM: Homo sapiens
160 <400> SEQUENCE: 2
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165             20                  25                  30
167 Gln Ile Ile Tyr Phe Asn Leu Glu Thr Val Gln Val Thr Trp Asn Ala
168         35                  40                  45
170 Ser Lys Tyr Ser Arg Thr Asn Leu Thr Phe His Tyr Arg Phe Asn Gly
171     50                  55                  60
173 Asp Glu Ala Tyr Asp Gln Cys Thr Asn Tyr Leu Leu Gln Glu Gly His
174 65                  70                  75                  80
176 Thr Ser Gly Cys Leu Leu Asp Ala Glu Gln Arg Asp Asp Ile Leu Tyr
177                 85                  90                  95
179 Phe Ser Ile Arg Asn Gly Thr His Pro Val Phe Thr Ala Ser Arg Trp
180             100                 105                 110
182 Met Val Tyr Tyr Leu Lys Pro Ser Ser Pro Lys His Val Arg Phe Ser
183         115                 120                 125
185 Trp His Gln Asp Ala Val Thr Val Thr Cys Ser Asp Leu Ser Tyr Gly
186     130                 135                 140
188 Asp Leu Leu Tyr Glu Val Gln Tyr Arg Ser Pro Phe Asp Thr Glu Trp
189 145                 150                 155                 160
191 Gln Ser Lys Gln Glu Asn Thr Cys Asn Val Thr Ile Glu Gly Leu Asp
192                 165                 170                 175
194 Ala Glu Lys Cys Tyr Ser Phe Trp Val Arg Val Lys Ala Met Glu Asp
195             180                 185                 190
197 Val Tyr Gly Pro Asp Thr Tyr Pro Ser Asp Trp Ser Glu Val Thr Cys
198         195                 200                 205
200 Trp Gln Arg Gly Glu Ile Arg Asp Ala Cys Ala Glu Thr Pro Thr Pro
201     210                 215                 220
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RAW SEQUENCE LISTING DATE: 03/05/2002
PATENT APPLICATION: US/10/078,059 TIME: 14:12:03

Input Set : A:\PF466P2SEQLIST01102002.txt
Output Set: N:\CRF3\03052002\J078059.raw

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203 Pro Lys Pro Lys Leu Ser Lys Phe Ile Leu Ile Ser Ser Leu Ala Ile
204 225                 230                 235                 240
206 Leu Leu Met Val Ser Leu Leu Leu Leu Ser Leu Trp Lys Leu Trp Arg
207                 245                 250                 255
209 Val Lys Lys Phe Leu Ile Pro Ser Val Pro Asp Pro Lys Ser Ile Phe
210             260                 265                 270
212 Pro Gly Leu Phe Glu Ile His Gln Gly Asn Phe Gln Glu Trp Ile Thr
213         275                 280                 285
215 Asp Thr Gln Asn Val Ala His Leu His Lys Met Ala Gly Ala Glu Gln
216     290                 295                 300
218 Glu Ser Gly Pro Glu Glu Pro Leu Val Val Gln Leu Ala Lys Thr Glu
219 305                 310                 315                 320
221 Ala Glu Ser Pro Arg Met Leu Asp Pro Gln Thr Glu Glu Lys Glu Ala
222                 325                 330                 335
224 Ser Gly Gly Ser Leu Gln Leu Pro His Gln Pro Leu Gln Gly Gly Asp
225             340                 345                 350
227 Val Val Thr Ile Gly Gly Phe Thr Phe Val Met Asn Asp Arg Ser Tyr
228         355                 360                 365
230 Val Ala Leu
231     370
235 <210> SEQ ID NO: 3
236 <211> LENGTH: 379
237 <212> TYPE: PRT
238 <213> ORGANISM: Homo sapiens
240 <400> SEQUENCE: 3
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242   1               5                  10                  15
244 Pro Leu Leu Gly Val Gly Leu Asn Pro Lys Phe Leu Thr Pro Ser Gly
245             20                  25                  30
247 Asn Glu Asp Ile Gly Gly Lys Pro Gly Thr Gly Gly Asp Phe Phe Leu
248         35                  40                  45
250 Thr Ser Thr Pro Ala Gly Thr Leu Asp Val Ser Thr Leu Pro Leu Pro
251     50                  55                  60
253 Lys Val Gln Cys Phe Val Phe Asn Val Glu Tyr Met Asn Cys Thr Trp
254 65                  70                  75                  80
256 Asn Ser Ser Ser Glu Pro Gln Pro Asn Asn Leu Thr Leu His Tyr Gly
257                 85                  90                  95
259 Tyr Arg Asn Phe Asn Gly Asp Asp Lys Leu Gln Glu Cys Gly His Tyr
260             100                 105                 110
262 Leu Phe Ser Glu Gly Ile Thr Ser Gly Cys Trp Phe Gly Lys Lys Glu
263         115                 120                 125
265 Ile Arg Leu Tyr Glu Thr Phe Val Val Gln Leu Gln Asp Pro Arg Glu
266     130                 135                 140
268 His Arg Lys Gln Pro Lys Gln Met Leu Lys Leu Gln Asp Leu Val Ile
269 145                 150                 155                 160
271 Pro Trp Ala Pro Glu Asn Leu Thr Leu Arg Asn Leu Ser Glu Phe Gln
272                 165                 170                 175
274 Leu Glu Leu Ser Trp Ser Asn Arg Tyr Leu Asp His Cys Leu Glu His
275             180                 185                 190
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RAW SEQUENCE LISTING DATE: 03/05/2002
PATENT APPLICATION: US/10/078,059 TIME: 14:12:03

Input Set : A:\PF466P2SEQLIST01102002.txt
Output Set: N:\CRF3\03052002\J078059.raw

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277 Leu Val Gln Tyr Arg Ser Asp Arg Asp Arg Ser Trp Thr Glu Gln Ser
278         195                 200                 205
280 Val Asp His Arg His Ser Phe Ser Leu Pro Ser Val Asp Ala Gln Lys
281     210                 215                 220
283 Leu Tyr Thr Phe Arg Val Arg Ser Arg Tyr Asn Pro Leu Cys Gly Ser
284 225                 230                 235                 240
286 Ala Gln His Trp Ser Asp Trp Ser Tyr Pro Ile His Trp Gly Ser Asn
287                 245                 250                 255
289 Thr Ser Lys Glu Asn Ile Glu Asn Pro Glu Asn Pro Ser Leu Phe Ala
290             260                 265                 270
292 Leu Glu Ala Val Leu Ile Pro Leu Gly Ser Met Gly Leu Ile Val Ser
293         275                 280                 285
295 Leu Ile Cys Val Tyr Cys Trp Leu Glu Arg Thr Met Pro Arg Ile Pro
296     290                 295                 300
298 Thr Leu Lys Asn Leu Glu Asp Leu Val Thr Glu Tyr Gln Gly Asn Phe
299 305                 310                 315                 320
301 Ser Ala Trp Ser Gly Val Ser Lys Gly Leu Ala Glu Ser Leu Gln Pro
302                 325                 330                 335
304 Asp Tyr Ser Glu Arg Leu Cys His Val Ser Glu Ile Pro Pro Lys Gly
305             340                 345                 350
307 Gly Glu Gly Pro Gly Gly Ser Pro Cys Ser Gln His Ser Pro Tyr Trp
308         355                 360                 365
310 Ala Pro Pro Cys Tyr Thr Leu Lys Pro Glu Pro
311     370                 375
314 <210> SEQ ID NO: 4
315 <211> LENGTH: 733
316 <212> TYPE: DNA
317 <213> ORGANISM: Homo sapiens
319 <400> SEQUENCE: 4
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322 tctcccggac tcctgaggtc acatgcgtgg tggtggacgt aagccacgaa gaccctgagg 180
323 tcaagttcaa ctggtacgtg gacggcgtgg aggtgcataa tgccaagaca aagccgcggg 240
324 aggagcagta caacagcacg taccgtgtgg tcagcgtcct caccgtcctg caccaggact 300
325 ggctgaatgg caaggagtac aagtgcaagg tctccaacaa agccctccca acccccatcg 360
326 agaaaaccat ctccaaagcc aaagggcagc cccgagaacc acaggtgtac accctgcccc 420
327 catcccggga tgagctgacc aagaaccagg tcagcctgac ctgcctggtc aaaggcttct 480
328 atccaagcga catcgccgtg gagtgggaga gcaatgggca gccggagaac aactacaaga 540
329 ccacgcctcc cgtgctggac tccgacggct ccttcttcct ctacagcaag ctcaccgtgg 600
330 acaagagcag gtggcagcag gggaacgtct tctcatgctc cgtgatgcat gaggctctgc 660
331 acaaccacta cacgcagaag agcctctccc tgtctccggg taaatgagtg cgacggccgc 720
332 gactctagag gat                                                   733
335 <210> SEQ ID NO: 5
336 <211> LENGTH: 5
337 <212> TYPE: PRT
338 <213> ORGANISM: Homo sapiens
340 <220> FEATURE:
341 <221> NAME/KEY: SITE
342 <222> LOCATION: (3)
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Use of n and / or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY DATE: 03/05/2002
PATENT APPLICATION: US/10/078,059 TIME: 14:12:04

Input Set : A:\PF466P2SEQLIST01102002.txt
Output Set: N:\CRF3\03052002\J078059.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:346 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:560 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:580 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:610 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:625 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:719 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:722 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:783 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:835 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:844 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:894 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30